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Human glucocorticoid receptor alpha mRNA,

PubMed, Protein, Related Sequences, Taxonomy, OMIM, LinkOut

complete cds

LOCUS HUMGCRA 4788 bp mRNA PRI 08-NOV-1994 DEFINITION Human glucocorticoid receptor alpha mRNA, complete cds.

ACCESSION M10901

VERSION M10901.1 GI:183032

KEYWORDS glucocorticoid receptor; glucocorticoid receptor-alpha.

SOURCE Human lymphoid cell line IM-9, cDNA to mRNA, clones hGR[1.2, 2.9,

5.16] and fibroblast cDNA library (H.Okayama), clones OB7 and OB10.

ORGANISM Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

REFERENCE (bases 1 to 4788)

**AUTHORS** Hollenberg, S.M., Weinberger, C., Ong, E.S., Cerelli, G., Oro, A.,

Lebo, R., Thompson, E.B., Rosenfeld, M.G. and Evans, R.M.

TITLE Primary structure and expression of a functional human

glucocorticoid receptor cDNA

**JOURNAL** Nature 318 (6047), 635-641 (1985)

MEDLINE 86092206

COMMENT Although [1] did not actually sequence this entire sequence from

one contiquous clone (thus eliminating the possibility that this sequence contains segments from multiple genes) their evidence strongly suggests that the alpha clone, OB7, and the beta clone, OB10, are transcribed from the same gene on chromosome 5 by alternate mRNA splicing. Therefore, the predicted alpha mRNA sequence is presented here in its entirety. Positions 1 to 648 were determined from clones OB10 and hGR5.16; positions 649-4788 were determined by overlapping regions of all 5 clones listed on the SOURCE line. The beta clone diverges from this alpha clone in sequence after position 2313. See also the beta GCR mRNA in entry with accession number M11050. [1] reports that the alpha form of glucocorticoid receptor is the predominant physiological form found in the various human and mouse cell lines that they tested. [1] also noted a region of chromosome 16 with enough homology to these clones to hybridize efficiently. Alternate polyadenylation signals present at positions 3101-3106 and 4678-4684 may also be utilized

by some mRNAs. A clone OB12 was isolated that used the 3101-3106

signal.

**FEATURES** Location/Oualifiers

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/map = "5q31 - q32"

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CDS 133..2466 /gene="GRL"

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http://www.../query.fcgi?cmd=Retrieve&db=Nucleotide&list\_uids=183032&dopt=GenBan \ 02/15/2001

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Nucleotide Protein Genome Structure PopSet Taxonomy OMIM Search Nucleotide Limits Index History Clipboard Default View as HTML SE)/6 Profesie Onsocient Human mRNA for beta-PubMed, Protein, Related Sequences, Taxonomy, OMIM, LinkOut glucocorticoid receptor (clone **OB10**) **HSGCRBR** LOCUS 3791 bp mRNA PRT 12-SEP-1993 DEFINITION Human mRNA for beta-glucocorticoid receptor (clone OB10). X03348 M11050 ACCESSION VERSION X03348.1 GI:31681 **KEYWORDS** glucocorticoid receptor. SOURCE human. ORGANISM Homo sapiens Eukaryota; Metazoa; Chordata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. REFERENCE (bases 1 to 3791) **AUTHORS** Hollenberg, S.M., Weinberger, C., Ong, E.S., Cerelli, G., Oro, A., Lebo, R., Thompson, E.B., Rosenfeld, M.G. and Evans, R.M. TITLE Primary structure and expression of a functional human qlucocorticoid receptor cDNA JOURNAL Nature 318 (6047), 635-641 (1985) MEDLINE 86092206 **FEATURES** Location/Qualifiers source 1..3791 /organism="Homo sapiens" /db xref="taxon:9606" CDS 133..2361 /note="(aa 1-742)" /codon start=1 /product="beta-glucocorticoid receptor" /protein id="CAA27054.1" /db xref="GI:31682" /db xref="SWISS-PROT:P04150" /translation="MDSKESLTPGREENPSSVLAQERGDVMDFYKTLRGGATVKVSAS SPSLAVASQSDSKQRRLLVDFPKGSVSNAQQPDLSKAVSLSMGLYMGETETKVMGNDL GFPQQGQISLSSGETDLKLLEESIANLNRSTSVPENPKSSASTAVSAAPTEKEFPKTH SDVSSEQQHLKGQTGTNGGNVKLYTTDQSTFDILQDLEFSSGSPGKETNESPWRSDLL IDENCLLSPLAGEDDSFLLEGNSNEDCKPLILPDTKPKIKDNGDLVLSSPSNVTLPQV KTEKEDF1ELCTPGV1KQEKLGTVYCQASFPGAN1IGNKMSA1SVHGVSTSGGQMYHY DMNTASLSQQQDQKPIFNVIPPIPVGSENWNRCQGSGDDNLTSLGTLNFPGRTVFSNG YSSPSMRPDVSSPPSSSSTATTGPPPKLCLVCSDEASGCHYGVLTCGSCKVFFKRAVE GQHNYLCAGRNDCIIDKIRRKNCPACRYRKCLQAGMNLEARKTKKKIKGIQQATTGVS QETSENPGNKT1VPATLPQLTPTLVSLLEV1EPEVLYAGYDSSVPDSTWR1MTTLNML GGRQVIAAVKWAKAIPGFRNLHLDDQMTLLQYSWMFLMAFALGWRSYRQSSANLLCFA PDLIINEQRMTLPCMYDQCKHMLYVSSELHRLQVSYEEYLCMKTLLLLSSVPKDGLKS QELFDEIRMTYIKELGKAIVKREGNSSQNWQRFYQLTKLLDSMHENVMWLKPESTSHT T.T " misc feature 3168..3173 /note="pot. polyA signal" 3539..3545 <u>misc</u> feature /note="pot. polyA signal" misc\_feature 3770..3775 /note="put. polyA signal" polyA site 3791 BASE COUNT 759 c 1162 a 808 g 1062 t ORIGIN 1 tttttagaaa aaaaaaatat atttccctcc tgctccttct gcgttcacaa gctaagttgt 61 ttatctcggc tgcggcggga actgcggacg gtggcgggcg agcggctcct ctgccagagt 121 tgatattcac tgatggactc caaagaatca ttaactcctg gtagagaaga aaaccccagc

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